

[illegible]**B**

1/6

FIG.2

A

Xenopus	1: ME-LLHTFCGGRWTL	UFTGILCFVARGV	GYYPFRSPFFFLCTHHGELEGDGEQGEVLI	59
mouse	1: RGCWAPRALVLAV	LLAT-RAF-AT	58
human	1: RSGWARQTFLLA	LGAT-RAF-AA	57
Xenopus	60: SLHLAGNPSYYTP	QGQEHVTISTSTFFDGLLV	TGLYSTSVQASQSIGGS	119
mouse	59:	118
human	58:	117
Xenopus	120: RQFGIQFMCSVVASHVSHLP	TITNLSFWIAP	PAGTGCNVFMATATHRGQVIFKDALAQQL	179
mouse	119: H.....	178
human	118: H.....	177
Xenopus	180: CEQGAPTEAPLRPN	AEIHSSEIL	LRDDFDSYKLQELNPNIWLQ	239
mouse	179:TAYSH.....	D.VI.....	QQL.....VE.S.....T.....	238
human	178:DVTVH.H.....	D.I.....	HQLQ.....VE.N.....T.....	237
Xenopus	240: GAVTFQDPYGPREDIT	VQNTITASVLQFSIGSGSCRF	SYSDPGIMVSYTKNNSSS	299
mouse	239: N.....E.....T.TCL.....	S.T.....A.....TAD.IQ.....	298
human	238: N.....E.....TGL.....	S.T.L.A.....AD.IQ.....	297
Xenopus	300: ERISAPSNVSTI	LHIIPLPEAKGNV	KRRNENMQADVYEACWALDNII	359
mouse	299: K.R.....V.....L.....E.....S.Q.Q.K	DSLRLV.E.....M.....S.....R.....	358
human	298: K.R.....L.....ED.....Q.Q.K.....LRV.E.....S.....RQ.....	357
Xenopus	360: VVLEDLDPMDTGNWLF	FPGATVKRILQSDGNSIYFHG	ESSEYRFATTROVDLSS	419
mouse	359:M.....S.....N.G.F.....T.....	418
human	358:S.....M.....S.....N.G.F.....T.....	417
Xenopus	420: DWSEEFENLPAQIV	GAVIGTECGTIESGSLVFLKDGERRV	CYPMOTTGYGNLRFY	479
mouse	419: E.....SQ.T.....D.L.....V.AD.....M.....L.....S.....	478
human	418: E.....SQ.T.....DVL.....M.....L.....S.....	477
Xenopus	480: FIMGGMLSPGESHEND	VLYAQIDGRKEHIIMOTLAFSSYKVT	SLVSAVTSFDLQTPATK	539
mouse	479: V.....I.D.M.....I.....K.E.....AL.....T.....P.....M.....N.E.....	538
human	478: V.....I.D.N.....I.....K.E.....TL.....S.....P.....M.....N.E.....	537
Xenopus	540: FCLKTSPRGHNLVWAVD	VHVLPLPSTESHMVQFSINLGGCS	HQPGNSVSLEFSTNH	599
mouse	539:R.K.....Q.Y.R.....F.....M.....I.....I.....	598
human	538:R.KN.Q.....R.....F.....M.....I.....I.....	597
Xenopus	600: GRSWCLTHSECLPELCAQSHLPHS
mouse	599:T.....I.....P.....
human	598:T.....I.....P.....

B

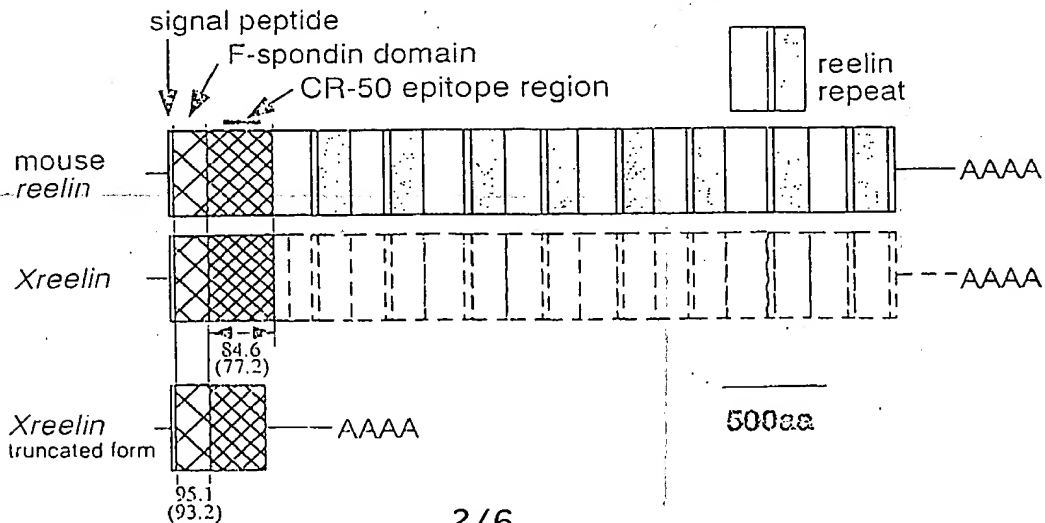


FIG.3

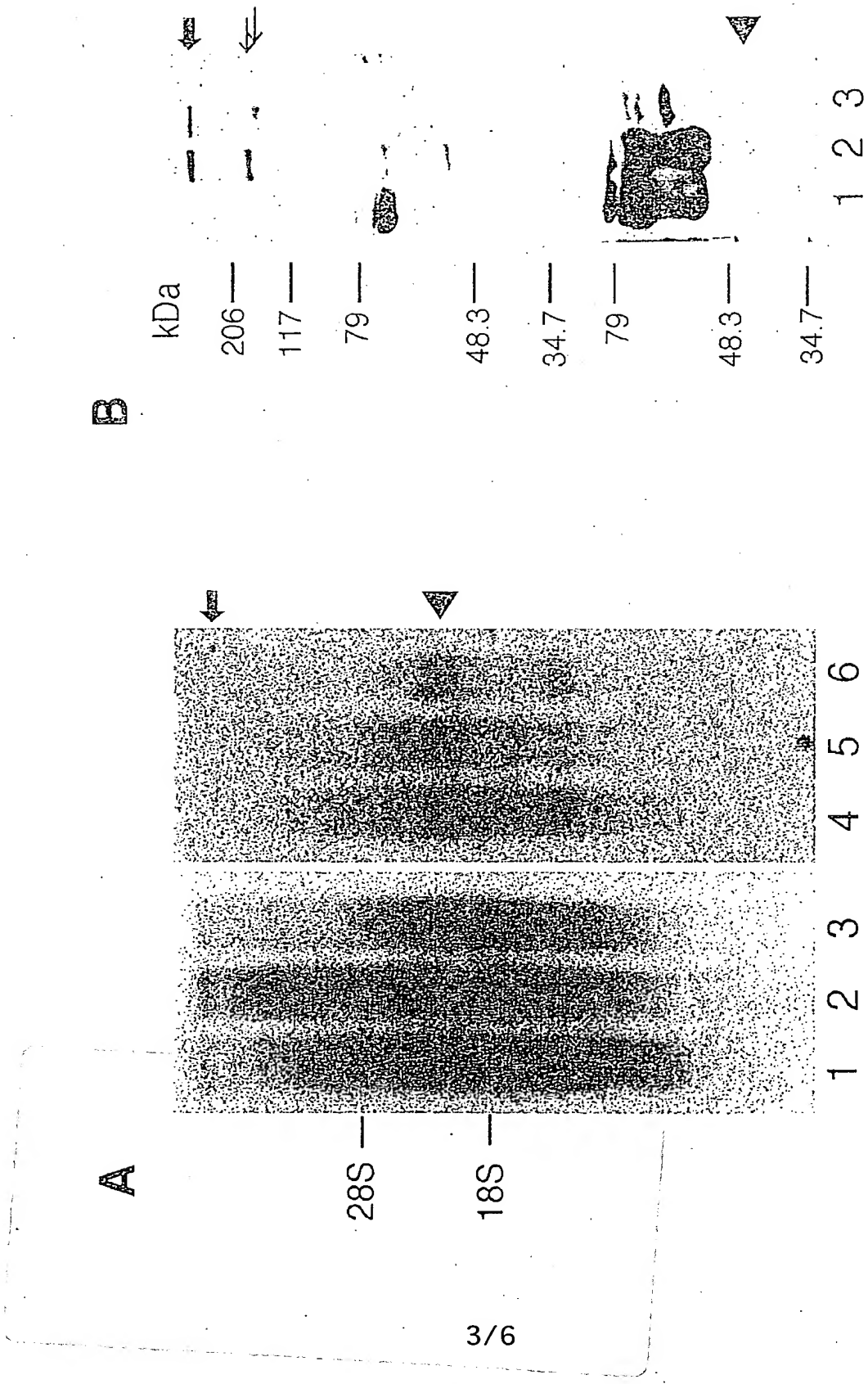


FIG.4

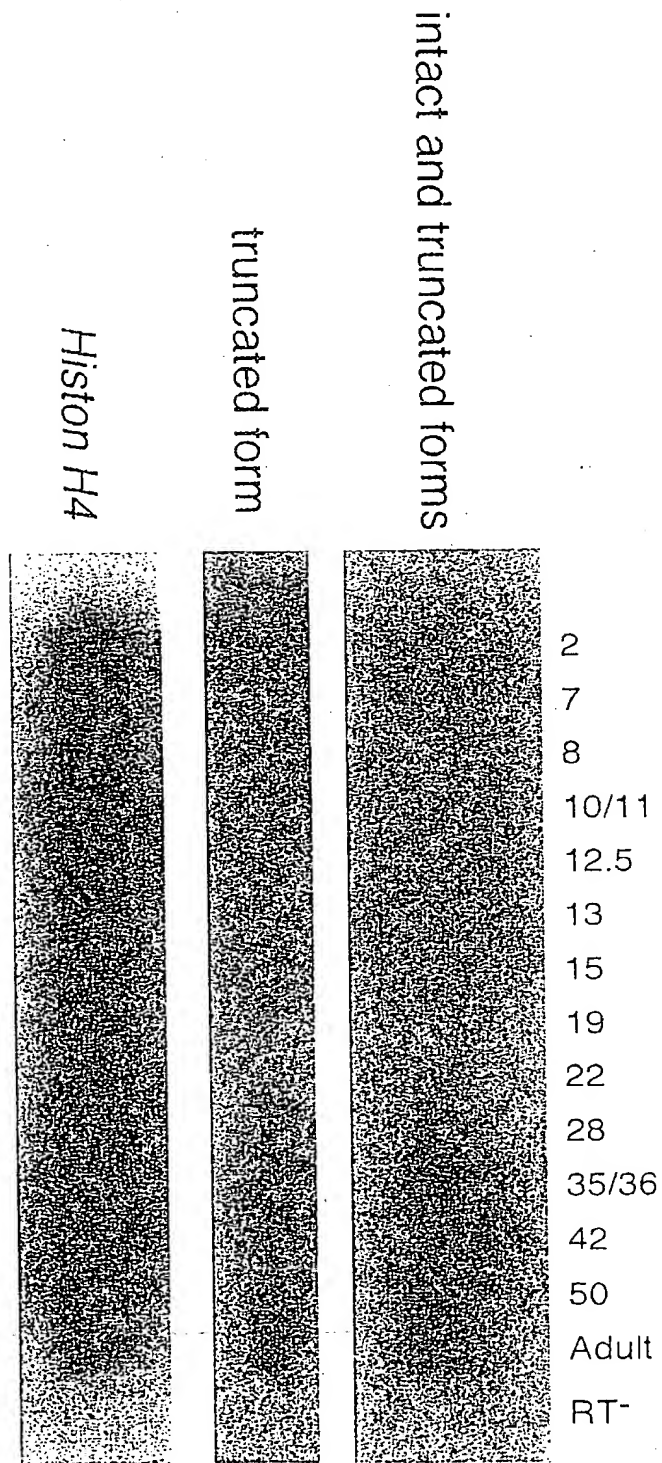


FIG.5

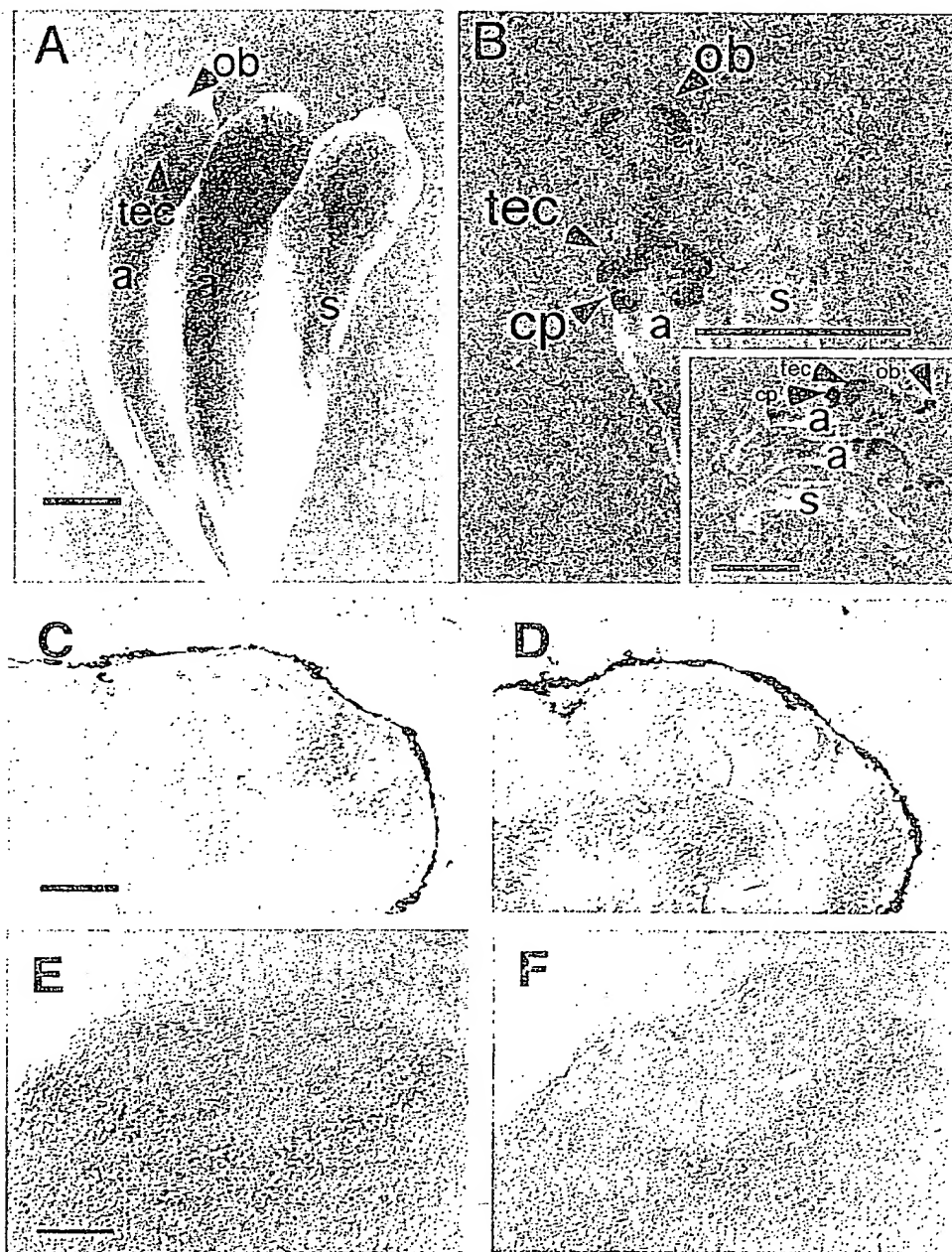


FIG.6

